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### Studies on the manipulation of gastrointestinal tract bacteria

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# **Studies on the Manipulation of Gastrointestinal Tract Bacteria**

*A thesis submitted in fulfillment of the requirements for the award of the degree*

**Master of Science (Research)**

from



by

**Peter Njuguna, MSc**

**School of Biological Sciences**

**2005**

## ABSTRACT

Increasing awareness that the human intestinal flora is a major factor in health and disease has led to different strategies to manipulate the flora to promote health. These approaches include changes to the diet by inclusion of prebiotics and probiotics. Prebiotics are non-digestible polysaccharide food ingredients that beneficially affect the host by selectively stimulating the growth and/or activity of one or a limited number of bacteria in the gastrointestinal tract (GIT). Probiotics on the other hand are viable culture of bacteria, which applied to animals or humans, beneficially affect the host by improving the properties of the indigenous microflora. One mechanism of action of probiotics is the production of antimicrobial substances called bacteriocins, such as colicins, that inhibit the growth of their competitors. The experiments described in this thesis examined the potential use of prebiotics and probiotics to manipulate GIT bacteria.

A crude polysaccharide extract (HW) from the medicinal mushroom *Ganoderma lucidum* was prepared by extracting the fruiting body with boiling water. The extract was then purified by ethanol precipitation resulting in the hot water-ethanol (HWE) extract. Groups of mice were fed these extracts over a period of three weeks at a concentration of 150 µg/ml in sterile drinking water and the mice then euthanised after three weeks. Changes in population dynamics of lumen bacteria were determined in the duodenum, ileum, colon and faeces by rigorous washing of excised segments while adherent bacteria were released with the non-ionic detergent Triton X100, which does not affect the viability of the bacteria. The prevalence of haemolytic colonies was assessed by plating washouts onto blood agar. Total colony forming units were enumerated on bacteriological media selective for *Enterobacteriaceae*, *Streptococci*, *Enterococci* and Lactic acid Bacteria (LAB). Results showed that there was little change in population dynamics elicited by extract feeding. The exception was a significant reduction in haemolytic lumen bacteria and increase in LAB lumen bacteria recovered from the colon of HWE treated mice.

A multiplex PCR was optimized and applied to survey the prevalence of eight common colicin genes (Colicins A, D, E1, E2, E6, E7, Ia and V) in *Escherichia coli* (*E. coli*) isolates. The study focused on 39 clinical isolates from

humans and 68 isolates from pigs with post-weaning diarrhoea. In addition, 152 porcine commensal *E. coli* isolates obtained from different compartments of the GIT (duodenum, ileum, colon and faeces) were included in the PCR analysis. Six individual colicins (E1, E2, E6, E7, Ia and V) and four dual colicin combinations (E1/E2, E1/E7, E2/E7, & E2/Ia) were detected. Approximately 28.2 % of the human pathogenic isolates had at least one colicin gene with colicins D, E1, E7 and V occurring at frequencies of 5.1 % each. Colicins E6, Ia and the dual colicin, E2/Ia, were less frequent and were found in about 2.6 % of clones. Only 4 % of the porcine pathogenic isolates possessed a colicin gene and these were exclusively E1 and V. In contrast, there was a significantly higher carriage (36.2%) of colicin genes in commensal porcine *E. coli*. Of these, E1, E7 and Ia accounted for 87 % of all colicin genes detected. Six of the commensal strains possessed multiple types of colicins with the most common being the E2/E7 combination. Furthermore, there appeared to be differences in the type of colicins found in commensal *E. coli* isolates recovered from different intestinal compartments.

Seven porcine commensal *E. coli* strains producing standard colicins were evaluated for inhibitory activity against five pathogenic *E. coli* of human and porcine origin. The experiment utilized a kinetic inhibitory microtitre assay (KIMA) to assess inhibition using non-induced supernatants and supernatants induced with 0.2 µg/ml of mitomycin C to stimulate colicin production. The level of inhibition was found to be variable with most of the commensal porcine *E. coli* strains showing little or no inhibitory action against the five pathogenic strains. However, two commensal strains, ILC33 and CC89 were found to be highly inhibitory to three porcine pathogenic *E. coli* strains of serotypes O141:K85, O141:K88 and O149:K88.

The findings of this thesis suggest that purified polysaccharide extracts (HWE) from *Ganoderma lucidum* have the potential to be used in further studies as prebiotics in view of their positive effects on beneficial LAB. In addition, the use of colicin-bearing strains as probiotic bacteria is justifiable because of the low incidence of colicin genes in pathogenic *E. coli* compared to commensals. Finally, these findings indicate that potential probiotic bacterial strains have to be scrutinised for their inhibitory activity against individual pathogenic strains prior to being subjected to further assessments.

## **CERTIFICATION**

I, Peter Njuguna, declare that this thesis, submitted in partial fulfilment of the requirements for the award of the degree of Master of Science (Research), in the School of Biological Sciences, University of Wollongong, is wholly my own work unless otherwise referenced or acknowledged. The document has not been submitted for qualification at any other academic institution.

Peter Njuguna

## **PUBLICATIONS AND PRESENTATIONS**

Njuguna, P., Wu, K., Chapman T., Chao, R., Zhang, R., Gordon, D., Bettelheim, K., and Chin, J. (2004). *E. coli* at War: Commensals versus Pathogens. PP12.3. Australian Society for Microbiology Conference, Sydney.

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# TABLE OF CONTENTS

<b>ABSTRACT</b>	<b>I</b>
<b>CERTIFICATE OF ORIGINALITY</b>	<b>III</b>
<b>PUBLICATIONS AND PRESENTATIONS</b>	<b>IV</b>
<b>ACKNOWLEDGEMENTS</b>	<b>V</b>
<b>TABLE OF CONTENTS</b>	<b>VI</b>
<b>LIST OF FIGURES</b>	<b>X</b>
<b>LIST OF TABLES</b>	<b>XII</b>
<b>LIST OF ABBREVIATIONS</b>	<b>XIII</b>
<b>Chapter 1: Review of Literature</b>	
<b>1.1 The Gastrointestinal Tract</b>	<b>1</b>
1.1.1 Beneficial Effects of Gastrointestinal Tract Microflora	<b>3</b>
1.1.2 Manipulation of Gastrointestinal Tract Microflora	<b>4</b>
1.1.3 Nutraceuticals	<b>5</b>
1.1.4 Prebiotics	<b>5</b>
1.1.5 Probiotics	<b>6</b>
<b>1.2 Medicinal Mushrooms</b>	<b>8</b>
1.2.1 General Information on Medicinal Mushrooms	<b>8</b>
1.2.2 Medicinal Use of Mushrooms	<b>9</b>
1.2.3 Genus <i>Ganoderma</i>	<b>10</b>
1.2.4 Polysaccharides in Medicinal Mushrooms	<b>11</b>
1.2.5 Polysaccharides with Immune Modulating Activity Isolated from Basidiomycetes	<b>12</b>
<b>1.3 Bacteriocins</b>	<b>15</b>
1.3.1 Colicins	<b>17</b>
1.3.2 Colicin Classification	<b>17</b>
1.3.3 Mechanism of Action	<b>18</b>
1.3.3.1 Nuclease Colicins	<b>19</b>
1.3.3.2 Channel-Forming Colicins	<b>21</b>
1.3.4 Colicin Plasmid Genes and their Products	<b>21</b>
1.3.4.1 Colicin Gene	<b>24</b>
1.3.4.2 Immunity Gene	<b>25</b>
1.3.4.3 Lysis Gene	<b>26</b>

1.3.5	Mechanism of Protection Against Channel-Forming and Nuclease Colicins	27
1.3.5.1	Resistance	28
1.3.5.2	Tolerance	29
1.3.6	Ecological Role of Colicin Production	30
<b>1.4</b>	<b>Aims of the Present Study</b>	<b>31</b>
 <b>Chapter 2: Effect of <i>Ganoderma lucidium</i> Fruiting Body Extracts on the Population Dynamics of Gastrointestinal Bacteria Grown on Selective Media</b>		
<b>2.1</b>	<b>Introduction</b>	<b>33</b>
<b>2.2</b>	<b>Materials and Methods</b>	<b>34</b>
2.2.1	Fruiting Body Extraction	34
2.2.1.2	Polysaccharide Content of Extracts	35
2.2.1.3	Protein Content of Extracts	36
2.2.2	Mice	36
2.2.2.1	Source	36
2.2.2.2	Holding Conditions	37
2.2.2.3	Feed	37
2.2.2.4	Care	37
2.2.2.5	Weight Estimations	38
2.2.3	Tissue Sampling	38
2.2.3.1	Anaesthesia	38
2.2.3.2	Removal of Intestinal Segments	38
2.2.3.3	Recovery of Luminal Bacteria / Digesta	39
2.2.3.4	Recovery of Adherent Bacteria	39
2.2.4	Faecal Sampling	40
2.2.4.1	Collection and Processing of Faeces	40
2.2.5	Bacteriological Media	40
2.2.5.1	Blood Agar	40
2.2.5.2	DeMan, Rogosa and Sharpe Agar	40
2.2.5.3	Kanamycin Esculin Azide Agar	41
2.2.5.4	Citric Azide Tween Carbonate Agar	41
2.2.5.5	MacConkey's (MAC) Agar	41
2.2.6	Processing and Enumeration of Bacteria	42
2.2.7	Experimental Protocol	42
2.2.8	Statistical Analysis	43
<b>2.3</b>	<b>Results</b>	<b>44</b>
2.3.1	Polysaccharide and Protein Content of Extracts	44
2.3.2	Weight Gain	44
2.3.3	Water Consumption	45
2.3.4	Bacterial Morphology on Selective Media	45

2.3.5	Haemolytic Colonies	48
2.3.6	Total Bacterial Enumeration	49
2.3.7	Lactic Acid bacterial Enumeration	52
2.3.8	<i>Streptococcal</i> Enumeration	53
2.3.9	<i>Enterococcal</i> Enumeration	56
2.3.10	<i>Enterobacteriaceae</i> Enumeration	58
<b>2.4</b>	<b>Discussion</b>	<b>61</b>
 <b>Chapter 3: Development of a Multiplex PCR Assay for the Characterisation of Colicinogenic Commensal <i>E. coli</i> Strains Inhibitory to Pathogenic <i>E. coli</i></b>		
<b>3.1</b>	<b>Introduction</b>	<b>65</b>
3.1.1	Aims	69
<b>3.2</b>	<b>Materials and Methods</b>	<b>70</b>
3.2.1	Bacterial and Colicinogenic Strains	70
3.2.2	Recovery of Bacteria and Preparation of Template DNA	74
3.2.3	Validation of Primer Pairs	75
3.2.3.1	PCR Amplification Conditions	75
3.2.3.2	Visualization of PCR Products by Gel Electrophoresis	76
3.2.4	Multiplex PCR	76
3.2.4.1	Reassessment of Multiplex PCR Conditions	78
3.2.5	Preparation of Colicin Supernatant Extracts from Reference and Producer Strains	79
3.2.6	Growth of Indicator Strains	80
3.2.7	Experimental Design for Assay	80
3.2.8	Calculations to Assess Effect of Bacterial Supernatant	81
<b>3.3</b>	<b>Results</b>	<b>83</b>
3.3.1	Validation of Primer Pairs	83
3.3.2	Multiplex PCR	85
3.3.3	Prevalence of Colicin Genes in Porcine Commensal <i>E. coli</i> Isolates	87
3.3.4	Prevalence of Colicin Genes in Different Intestinal Compartments	89
3.3.5	Frequency of Colicin Genes in Porcine and Human Pathogenic <i>E. coli</i> Isolates	92
3.3.6	Inhibitory Ability of Colicin Reference Strains on Colicin Sensitive Strain BZB1011	94
3.3.7	Growth Effect of Colicin Reference Strains on Selected <i>E. coli</i> Pathogens	97

3.3.8	Assessment of Effect of Commensal Strains on Colicin Sensitive Strain BZB1011	104
3.3.9	Assessment of Inhibitory Effect of Commensal Strains on Selected Pathogenic Strains	106
<b>3.4</b>	<b>Discussion</b>	<b>109</b>
<b>Chapter 4:</b>	<b>Conclusion and Future Directions</b>	<b>116</b>
<b>References</b>		<b>118</b>
<b>Appendix 1:</b>	<b>Bacteriological Media</b>	<b>128</b>
<b>Appendix 2:</b>	<b>Buffers and Solutions</b>	<b>130</b>
<b>Appendix 3:</b>	<b><i>E. coli</i> Isolates Screened for Colicin Genes</b>	<b>131</b>
<b>Appendix 4:</b>	<b>Volumes of PCR Cocktails Used for Optimised Multiplex PCR</b>	<b>137</b>

## LIST OF FIGURES

### Chapter 1

Figure 1.1	The human gastrointestinal tract	1
Figure 1.2.	Structure of branched (1-3)- $\beta$ -D-glucans	14
Figure 1.3.	Organization of the colicin-immunity-lysis gene operon	23
Figure 1.4.	Regions of the colicin molecule	25

### Chapter 2

Figure 2.1.	Scan of HW and HWE extracts from <i>Ganoderma lucidium</i>	44
Figure 2.2.	Changes in body weight over time due to mushroom extract feeding	46
Figure 2.3.	Average amount of water consumed per mouse per day in each treatment group	46
Figure 2.4.	Morphology of bacterial populations on different selective media	47
Figure 2.5.	Bacterial morphology on blood agar	47
Figure 2.6.	Effect of mushroom extracts feeding on the frequency of haemolytic colonies on blood agar	50
Figure 2.7.	Effect of mushroom extracts feeding on total bacterial enumeration on blood agar	51
Figure 2.8.	Effect of mushroom extracts feeding on Lactic acid bacterial enumeration on MRS agar	54
Figure 2.9.	Effect of mushroom extracts feeding on <i>Streptococcal</i> enumeration on KEA agar	55
Figure 2.10.	Effect of mushroom extracts feeding on <i>Enterococcal</i> enumeration on CATC agar	57
Figure 2.11.	Effect of mushroom extracts feeding on <i>Enterbacteriaceae</i> enumeration on MAC agar	60

### Chapter 3

Figure 3.1.	Longitudinal Cross-section of a 96-well microtitre plate	81
Figure 3.2	Agarose gel showing validation of primer pairs	84
Figure 3.3.	Agarose gel of initial amplification products obtained by multiplex PCR	86
Figure 3.4.	Agarose gel of multiplex PCR amplicons using final optimal conditions	87
Figure 3.5.	Frequency of colicin bearing commensal clones from each gastrointestinal compartment	88
Figure 3.6.	Frequency of colicin genes in commensal <i>E. coli</i> isolates from the duodenum and ileum	90
Figure 3.7	Distribution of colicin genes in commensal <i>E. coli</i> isolates from the colon and faeces	91
Figure 3.8	Frequency of colicin genes in pathogenic <i>E. coli</i> isolates from porcine and human sources	93

Figure 3.9.	Effects of colicin supernatant from growth cultures of <i>E. coli</i> reference strains on growth of colicin sensitive strain BZB1011	<b>95</b>
Figure 3.10.	Assessment of inhibition of selected pathogenic strains by <i>E. coli</i> reference strains	<b>99</b>
Figure 3.11	Assessment of inhibition of colicin sensitive strain BZB1011 by <i>E. coli</i> commensal strains	<b>105</b>
Figure 3.12	Assessment of inhibition of selected pathogenic strains by <i>E. coli</i> commensal strains	<b>107</b>

## LIST OF TABLES

### Chapter 1

Table 1.1 Characteristics of previously identified colicins	20
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### Chapter 3

Table 3.1. List of <i>E. coli</i> strains carrying plasmids encoding specific colicin genes	72
Table 3.2. Primer pairs used for detection of colicin genes by PCR	73
Table 3.3. Primer pools used for multiplex PCR	77
Table 3.4. Frequency and presence of each colicin type in different GIT compartments of porcine commensal <i>E. coli</i>	92

## LIST OF ABBREVIATIONS

BA	Blood agar
BP	Base pairs
CATC	Citric Azide Tween Carbonate agar
CC	Colon commensal
CFU	Colony forming units
DC	Duodenum commensal
<i>E. coli</i>	<i>Escherichia coli</i>
EHEC	Enterohemorrhagic <i>E. coli</i>
ETEC	Enterotoxigenic <i>E. coli</i>
FC	Faecal commensal
GIT	Gastrointestinal tract
HW	Hot water extract
HWE	Hot water-ethanol extract
ILC	Ileum commensal
KDa	Kilodaltons
KEA	Kanamycin Esculin Azide agar
LA	Luria agar
LAB	Lactic acid bacteria
LB	Luria agar broth
MAC	MacConkeys agar
ML	Milliliter
MRS	Deman, Rogosa and Sharpe agar
NDGOS	Non-digestible galacto-oligosaccharides
O.D.	Optical density
PBS	Phosphate Buffer Saline
PCR	Polymerase chain reaction
PWD	Post-weaning diarrhoea
RPM	Rotations per minute
SCFA	Short-chain fatty acids
µg	Micro-gram
TX	Triton-X 100